

**AMENDMENT****In the Specification:**

**Please add the below subheading and replace the paragraph under the title, on page 1, lines 4-8, with the following:**

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**Cross-Reference to Related Applications**

B1 This application is a continuation-in-part application of U.S. Ser. No. 09/323,597 filed June 1, 1999 which claims the benefit of United States provisional patent application serial numbers 60/087,598, filed June 1, 1998, now lapsed, 60/091,474 filed June 29, 1998, now lapsed and 60/129,521, filed April 14, 1999, now lapsed. The entire contents of these provisional and non-provisional patent applications are incorporated by reference into this application.

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**Please replace the paragraph from page 16, line 32 through page 17, line 7 with the following:**

B2 In the context of amino acid sequence comparisons, the term "identity" is used to express the percentage of amino acid residues at the same relative positions that are the same. Also in this context, the term "homology" is used to express the percentage of amino acid residues at the same relative positions that are either identical or are similar, using the conserved amino acid criteria of BLAST analysis, as is generally understood in the art. For example, % identity values may be generated by WU-BLAST-2 (Altschul et al., 1996, Methods in Enzymology 266:460-480; available at http address "blast.wustl/edu/blast/README.html"). Further details regarding amino acid substitutions, which are considered conservative under such criteria, are provided below.

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